

Figure 4C

		10	20	30	40	. 50	60
		.		1	1	• • • • • • •	
Table 6	MGQCRSAN	AEDAQEF	SDVERALET	LIKNFHQYSV	EGGKETLTPSI	ELRDLVTQQ	LPHLMPS
AY007220	MGQCRSAN	AEDAQEF	SDVERAIET	LIKNFHQYSV	EGGKETLTPSI	ELRDLVTQQ	LPHLMPS
Consensus	MGQCRSAN	AEDAQEF	SDVERAIET	LIKNFHQYSV	EGGKETLTPS	ELRDLVTQQ	LPHLMPS
		70	80	90	100		
		.		[l	1		
Table 6	NCGLEEKI	ANLGSCN	DSKLEFRSF	WELIGEAAKS	VKLERPVRGH		
AY007220 Consensus					VKLERPVRGH		NO:39)
	NCGLEEKI	ANLGSCN	DSKLEFRSF	WELIGEAAKS	VKLERPVRGH.	(SEQ ID	NO:40)

Figure 4D

30 (SEQ ID NO:3) Table 3 (SEQ ID NO:41) gi | 4139958 | pdb | 1MHO (SEQ ID NO:42) FHOYSRREGDKOTLTE PROTEIN MRP-126 YS<mark>GKEGD</mark>K (SEQ ID NO:43) CMALLISTEH ICTACALCIN (SEQ ID NO:44) CALGRANULIN B (SEQ ID NO:45) Consensus

Figure 4E

Table 6
gi|4139958|pdb|1MHO
PROTEIN MRP-126
CALGRANULIN B
CALGRANULIN B
Consensus

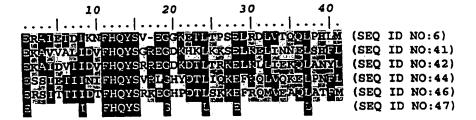






Figure 7

65677221+	GAATTCCAGAGGAGTTCTCAGTGCCCCCGGACAGGCCTCTCCAGCTTCACACTCTTGGC
AA315020-	TGCCCCGGACAGTCCTCTCNAGCTTCACACTCTTGGC
consensus	GAATTCCAGAGGAGTTCTCAGTGCCCCCGGACAGGCCTCTCCAGCTTCACACTCTTGGC
65677221+	CGCTTCTCCAATCAGCTCCCAGAAACTCCTGAACTCCAGTTTAGAGTCATTGCAGCTGCC
AA315020-	CGCTTCTCCAATCAGCTCCCAGAAACTCCTGAACTCCAGTTTAGAGTCATTGCAGCTGCC
consensus	CGCTTCTCCAATCAGCTCCCAGAAACTCCTGAACTCCAGTTTAGAGTCATTGCAGCTGCC
65677221+	CAGGTTGGCAATTTTCTCTTCCAGGCCACAGTTGCTCGGCATGAGATGGGGCCAGCTGCTG
AA315020-	CAGGTTGGCAATTTTCTCTTCCAGGCCANAGTTGCTCGGCATGAGATGGGGCCAGCTGCTG
consensus	CAGGTTGGCAATTTTCTTCTTCCAGGCCACAGTTGCTCGGCATGAGATGGGGCCAGCTGCTG
65677221+	GGTGACCAGGTCCCGTAGCTCAGAAGGGGTCAGCGTCTCCTTCCCACCGCACGAGTA
AA315020-	GGTGACCAGGTCCCGTAGCTCAGAGGGGTCAGCGTCTCCCTTCCCACGGAGTA
consensus	GGTGACCAGGTCCCGTAGCTCAGAAGGGGTCAGCGTCTCCCTTCCCACGGAGTA
65677221+	CTGGTGAAAGTTCTTGATGAGGGTCTCAATGGCCCTCTCCACATCACTGAATTC (SEQ ID NO: 37)
AA315020-	CTGGTGAAAGTTCTTGATGAGGGTCTCAATGGCCCTCTCCACATCACTGAATTCCTGAGC
consensus	CTGGTGAAAGTTCTTGATGAGGGTCTCAATGGCCCTCTCCACATCACTGAATTCCTGAGC
AA315020-	ATCCTCTGCGTTGGCTGACCGACACTGTCCCATGGTGCTCACTGTGTCTGGTCCTTTGGT
consensus	ATCCTCTGCGTTGGCTGACCGACACTGTCCCATGGTGCTCACTGTGTCTGGTCCTTTGGT
AA315020-	GAGAGTTCTGTTGTCCTAT (SEQ ID NO: 49)
consensus	GAGAGTTCTGTTGTCCTAT (SEQ ID NO: 5)